

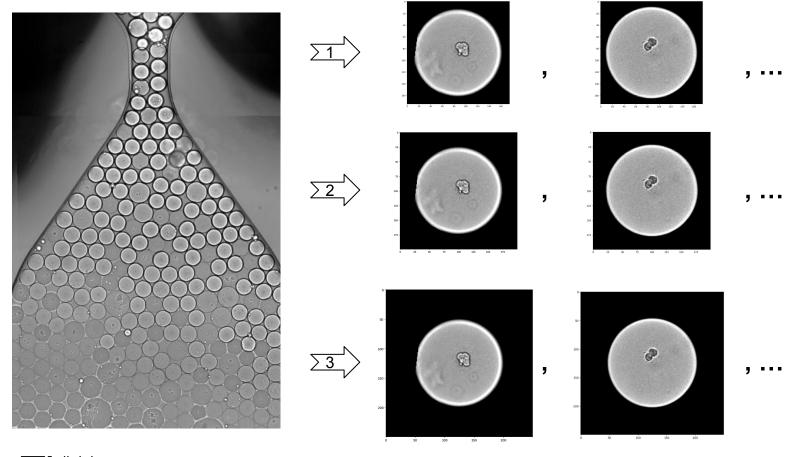


Our role as data scientists

- Find a way to automate the classification of tumor droplets.
- Use deep learning to achieve this.
- Make use of manually annotated tumor droplets for training.
- Draw conclusions about response of different drugs.
- Jupyter notebook as interface to use our code



Data Preprocessing



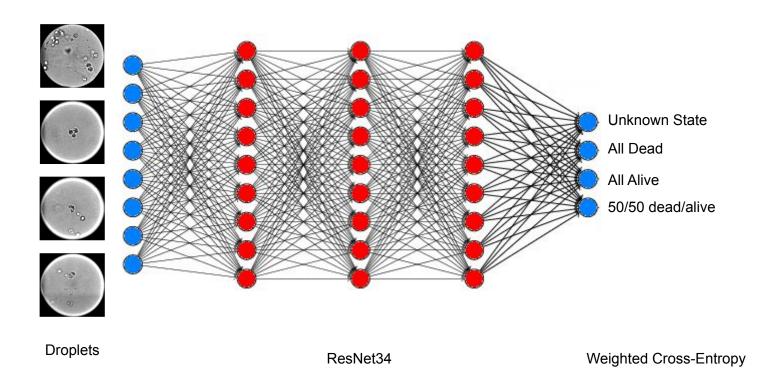
ETH zürich

Always use the same preprocessing for training and prediction!

Demonstration of preprocessing in the Notebook

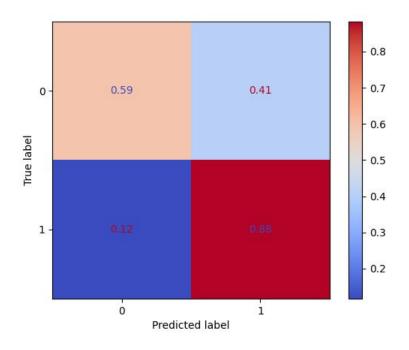


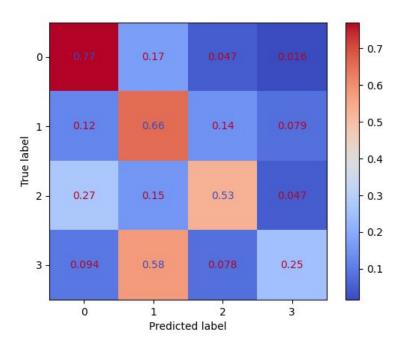
Model & Training





Results







Demonstration of training in the Notebook

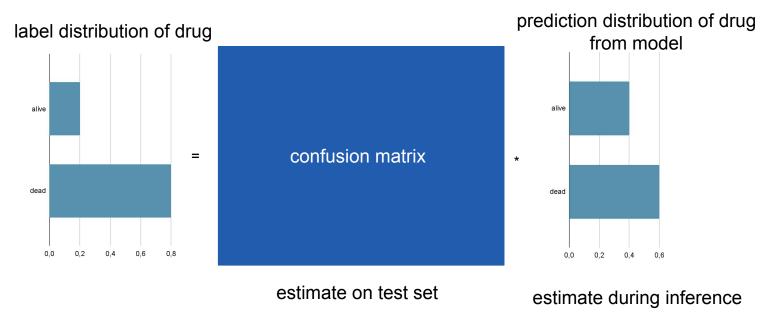


Key insight: We only care about *overall* drug effect per patient



Estimating Label Distribution per Drug

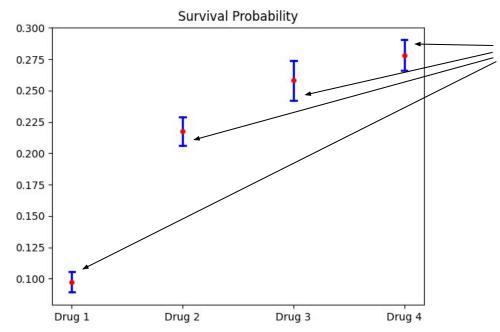
With the right statistical model:



How good are the estimates?



Different Drugs, Different Survival



 $(1-\alpha)$ - confidence intervals for class alive:

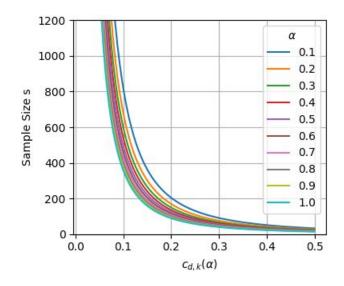
- 1. Derived from concentration inequalities.
- 2. Hold uniform over the drugs.
- If they do not overlap:
 With probability (1-α) we have found the best drug!

Note: this plot is a sketch and was not computed using actual data.



More samples means higher confidence

Deviation vs. sample size for different confidences for C = 2





A word of caution

- These confidence bounds depend on mathematical assumptions.
- If those assumptions do not hold, you cannot trust the results.
- Main advantage of this approach:
 We can get some idea of the uncertainty in or

We can get some idea of the uncertainty in our estimates.



Demonstration of prediction in the Notebook



Thank you! Any questions?

